

# Infectious Disease Models 1

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CMPT 858

March 9, 2010

# Comments on Mathematics & Dynamic Modeling

- Many accomplished & well-published dynamic modelers have very limited mathematical background
  - Can investigate pressing & important issues
  - Software tools are making this easier over time
- Can gain extra insight/flexibility if willing to push to learn some of the associated mathematics
- Achieving highest skill levels in dynamic modeling do require mathematical facility and sophistication
  - To do sophisticated work, often those lacking this background or inclination collaborate with someone with background

# Applied Math & Dynamic Modeling

- Although you may not use it, the dynamic modeling presented rests on the tremendous deep & rich foundation of applied mathematics
  - Linear algebra
  - Calculus (Differentia/Integral, Uni& Multivariate)
  - Differential equations
  - Numerical analysis (including numerical integration, parameter estimation)
  - Control theory
  - Real & complex analysis
- For the mathematically inclined, the tools of these areas of applied math are available

# Models in Mathematical Epidemiology of Infectious Disease

- Long & influential modeling tradition, beginning with Kermack-McKendrick (1920s)
- Models formulated for diverse situations (Cf. Anderson & May)
  - Latent & incubation period/Diversity in contact rates/Heterogeneity/Preferential mixing/Vaccinated groups/Zoonoses/Variations in clinical manifestations/Network structure
- Important tradition of closed-form analysis

# Mathematical Models Link Together Diverse Factors

## *Typical Factors Included*

- Infection
  - Mixing & Transmission
  - Development & loss of immunity – both individual and collective
  - Natural history (often multi-stage progression )
  - Recovery
- Birth & Migration
- Aging & Mortality
- Intervention impact

## *Sometimes Included*

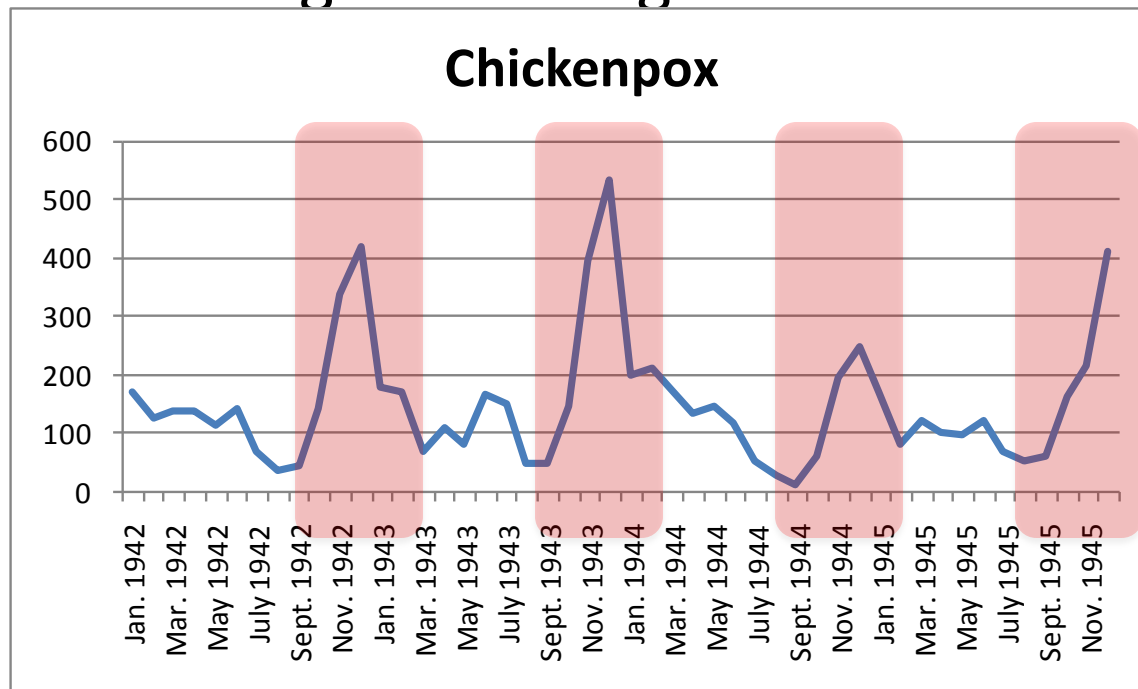
- Preferential mixing
- Variability in contacts
- Strain competition & cross-immunity
- Quality of life change
- Health services interaction
- Local perception
- Changes in behavior, attitude
- Immune response

# Emergent Characteristics of Infectious Diseases Models

- Instability
- Nonlinearity
- Tipping points
- Oscillations
- Multiple fixed points/equilibria
  - “Endemic” equilibrium
  - Disease free equilibrium

# Instability

- Slight perturbation (e.g. arrival of infectious person on a plane) can cause big change in results
  - Contrast with “goal seeking” behaviour

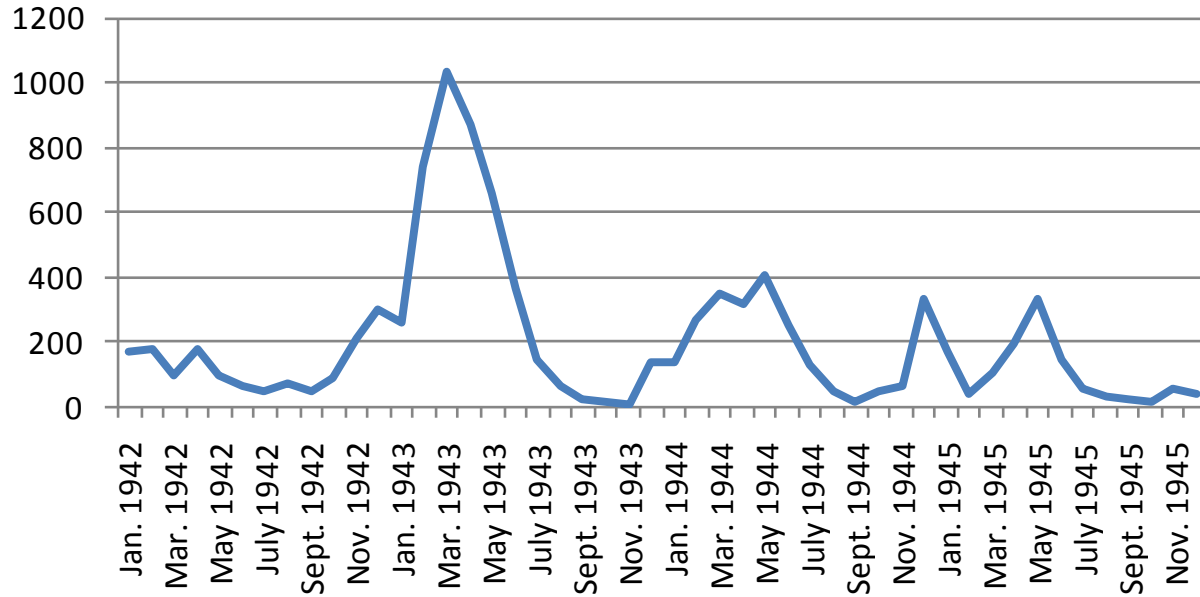


# Oscillations & Delays

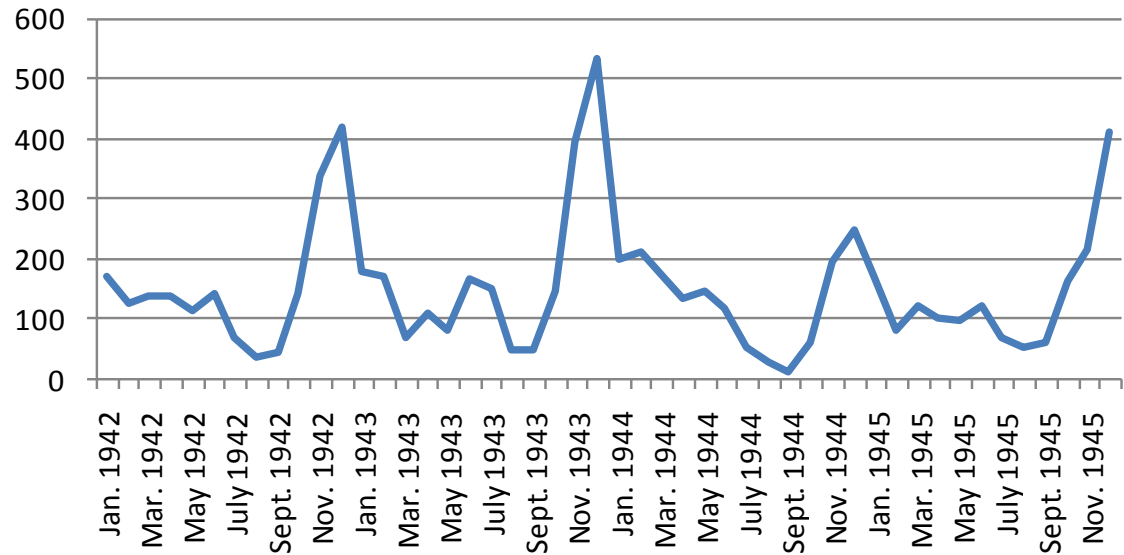
- The oscillations reflect negative feedback loops with delays
- These delays reflect “stock and flow” considerations and specific thresholds dictating whether net flow is positive or negative
  - Stock & Flow: Stock continues to deplete as long as outflow exceeds inflow, rise as  $\text{inflow} > \text{outflow}$ 
    - The stock may stay reasonably high long after inflow is low!
  - Key threshold  $R^*$ : When # of individuals being infected by a single infective = 1
    - This is the threshold at which  $\text{outflows} = \text{inflows}$



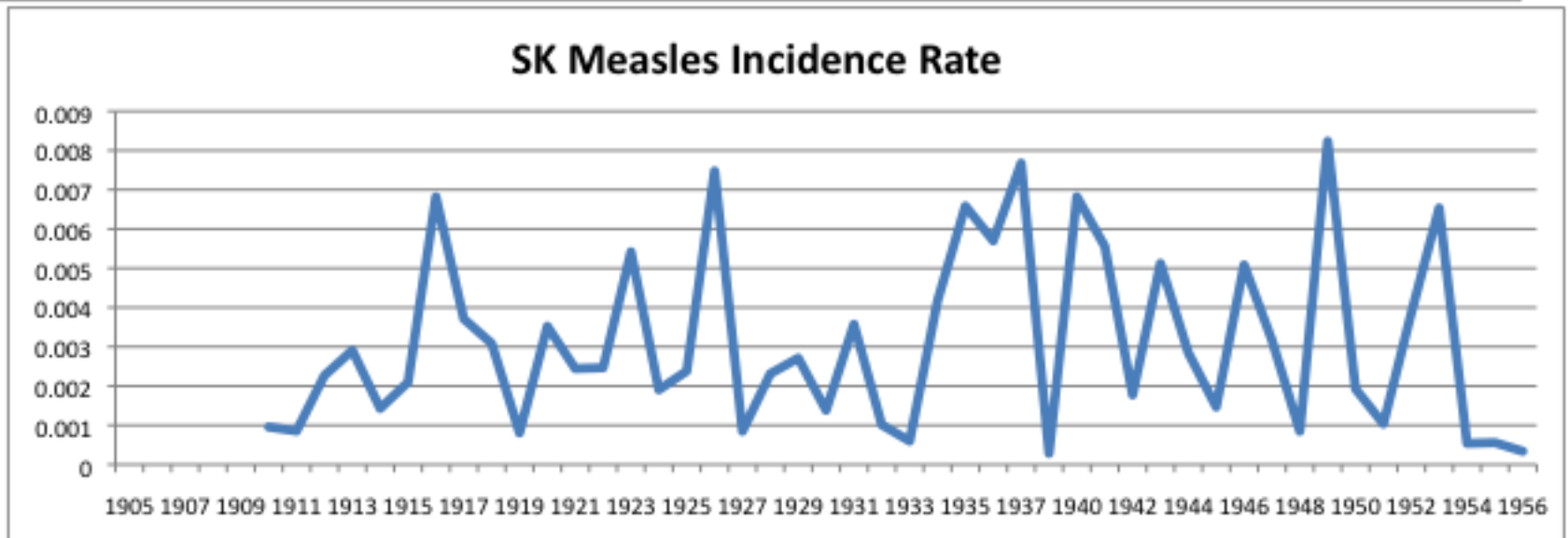
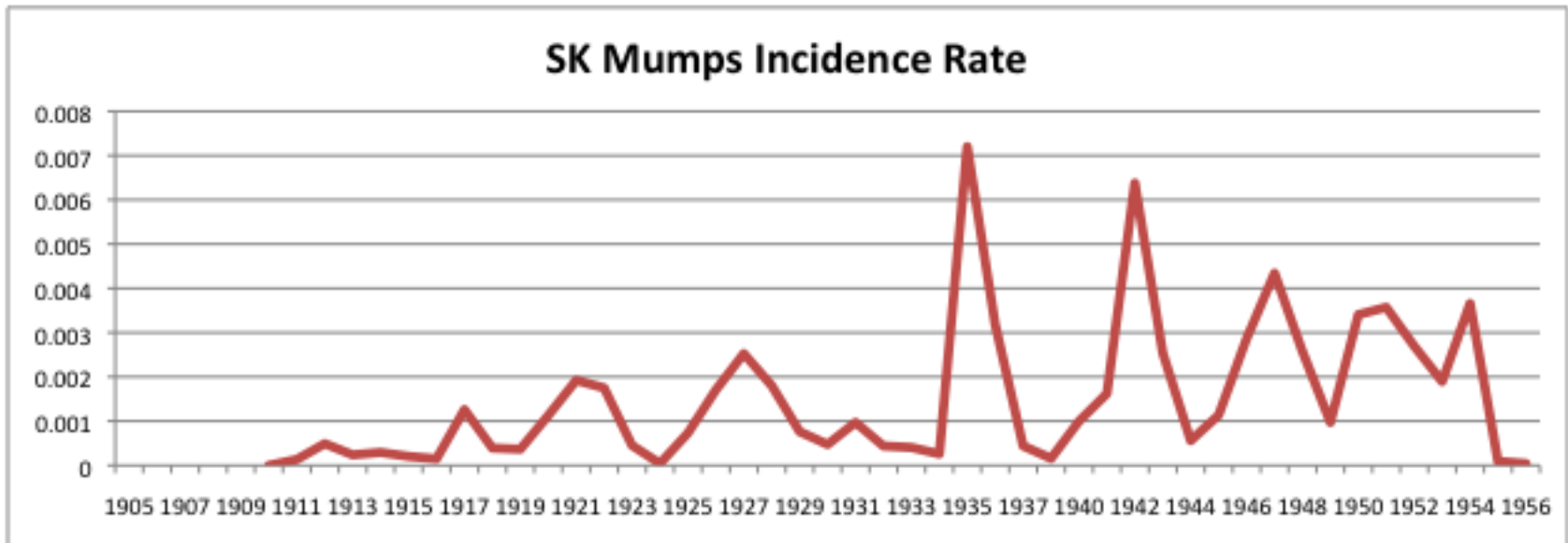
# Measles



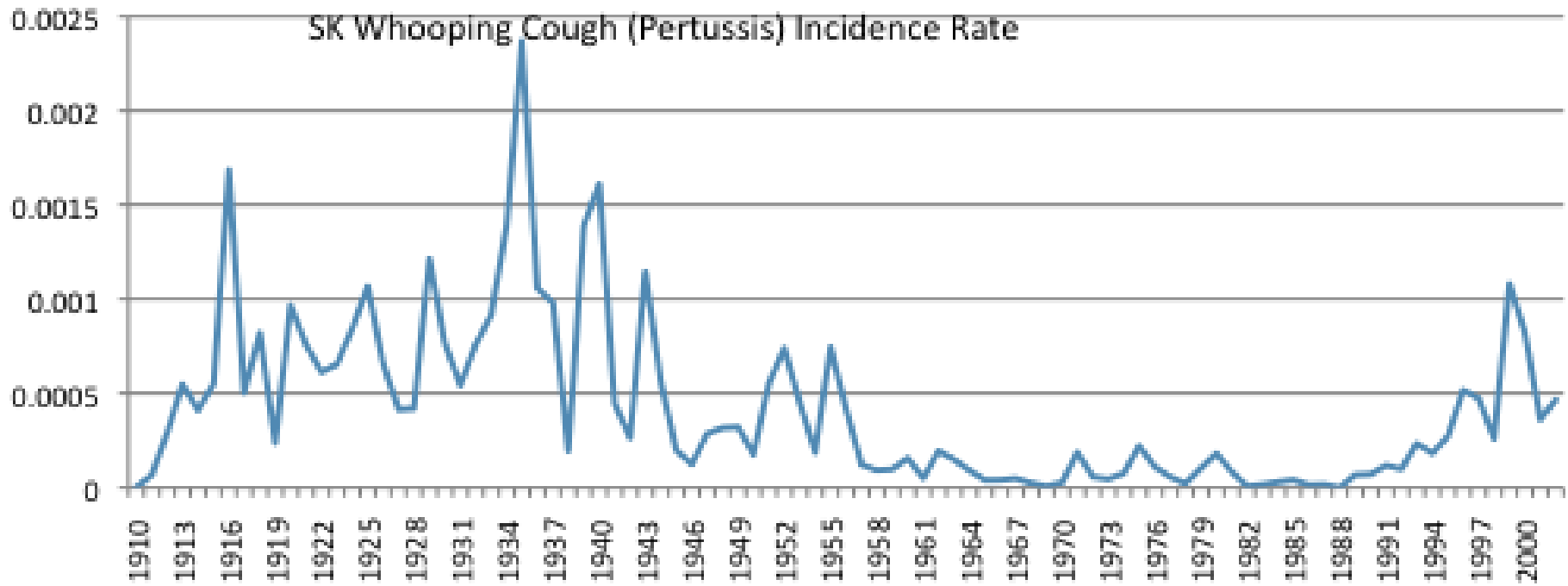
# Chickenpox



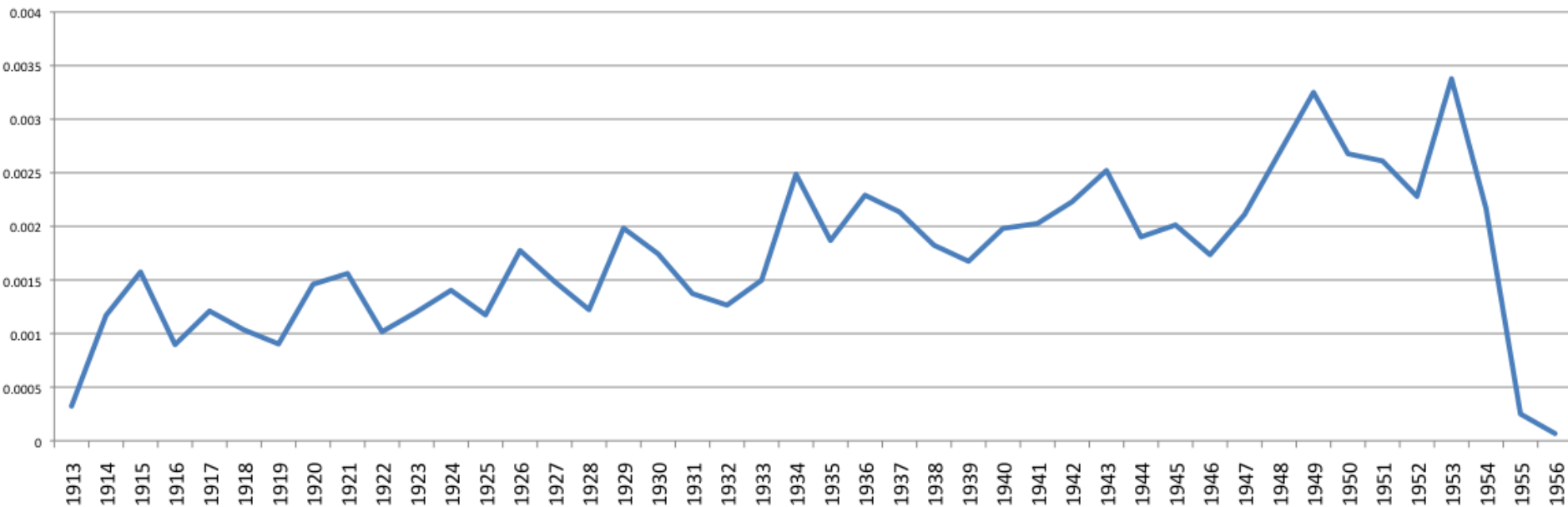
# Saskatchewan Childhood Diseases



### SK Whooping Cough (Pertussis) Incidence Rate



### SK Chicken Pox Incidence Rate



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Reasons

# Nonlinearity (in state variables)

- Effect of multiple policies non-additive
- Doubling investment does not yield doubling of results
- Leads to
  - Multiple basins of tracking (equilibrium)

# Multiple Equilibria & Tipping Points

- Separate basins of attraction have qualitatively different behaviour
  - Oscillations
  - Endemic equilibrium
  - Disease-free equilibrium

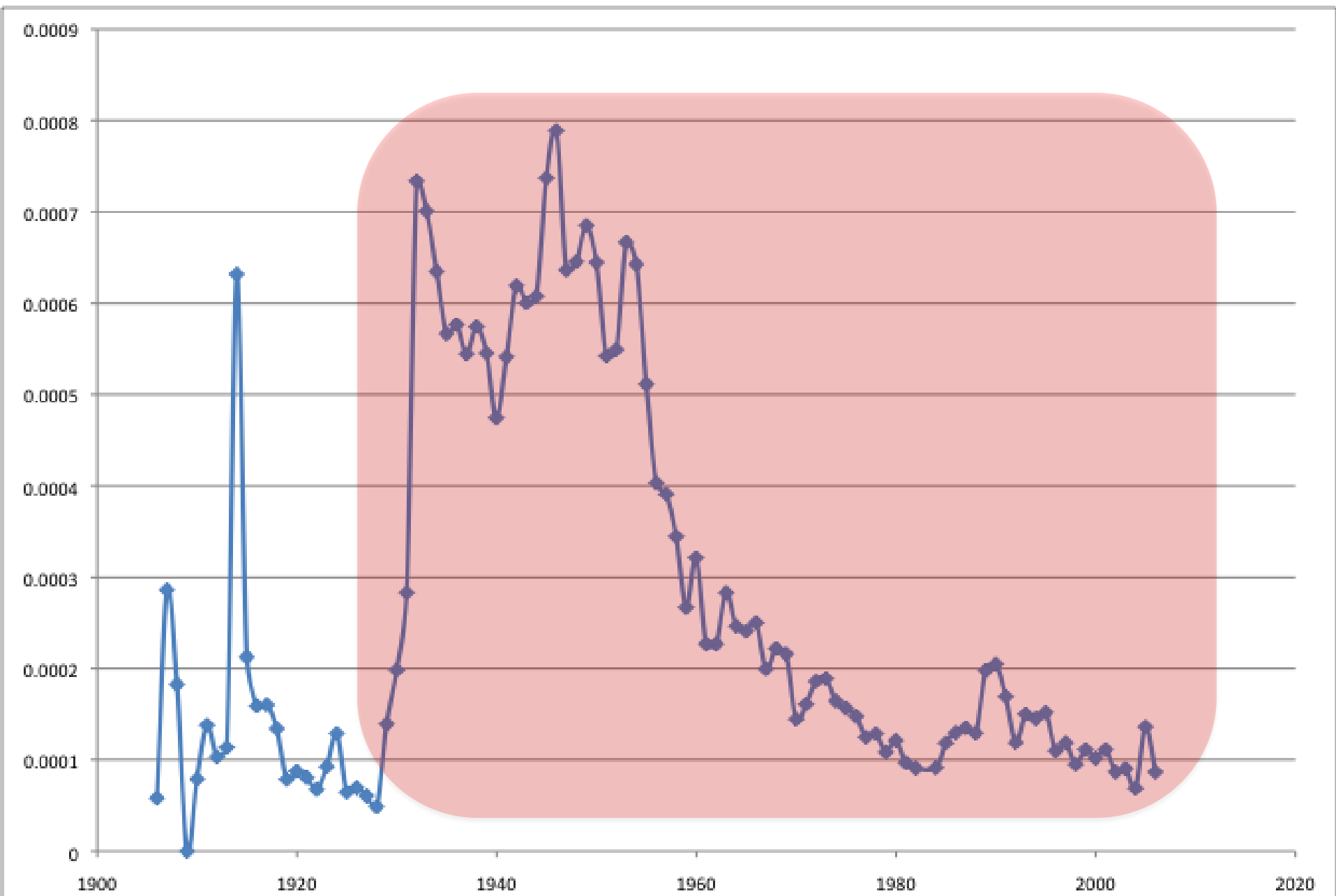
# Equilibria

- Disease free
  - No infectives in population
  - Entire population is susceptible
- Endemic
  - Steady-state equilibrium produced by spread of illness
  - Assumption is often that children get exposed when young

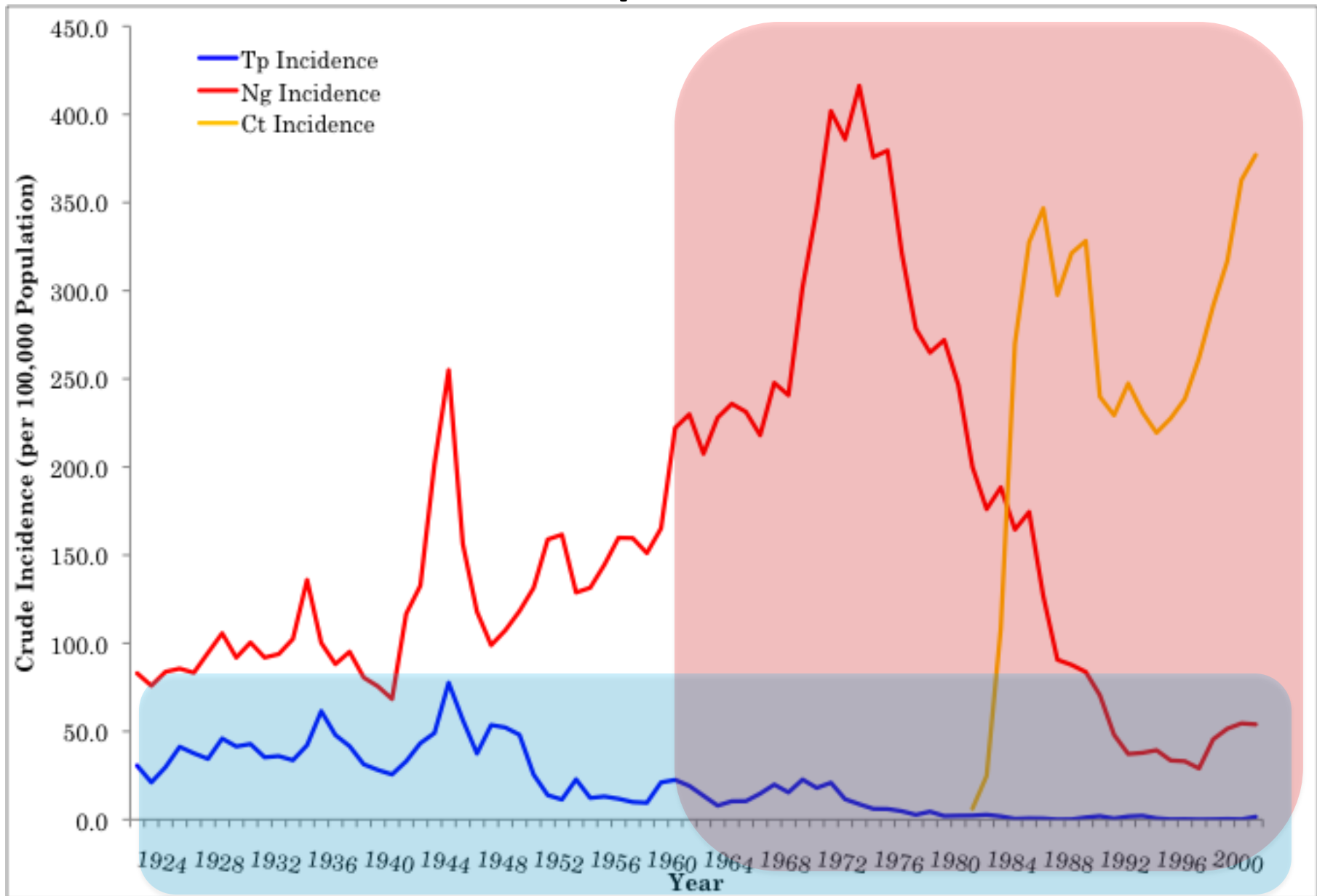
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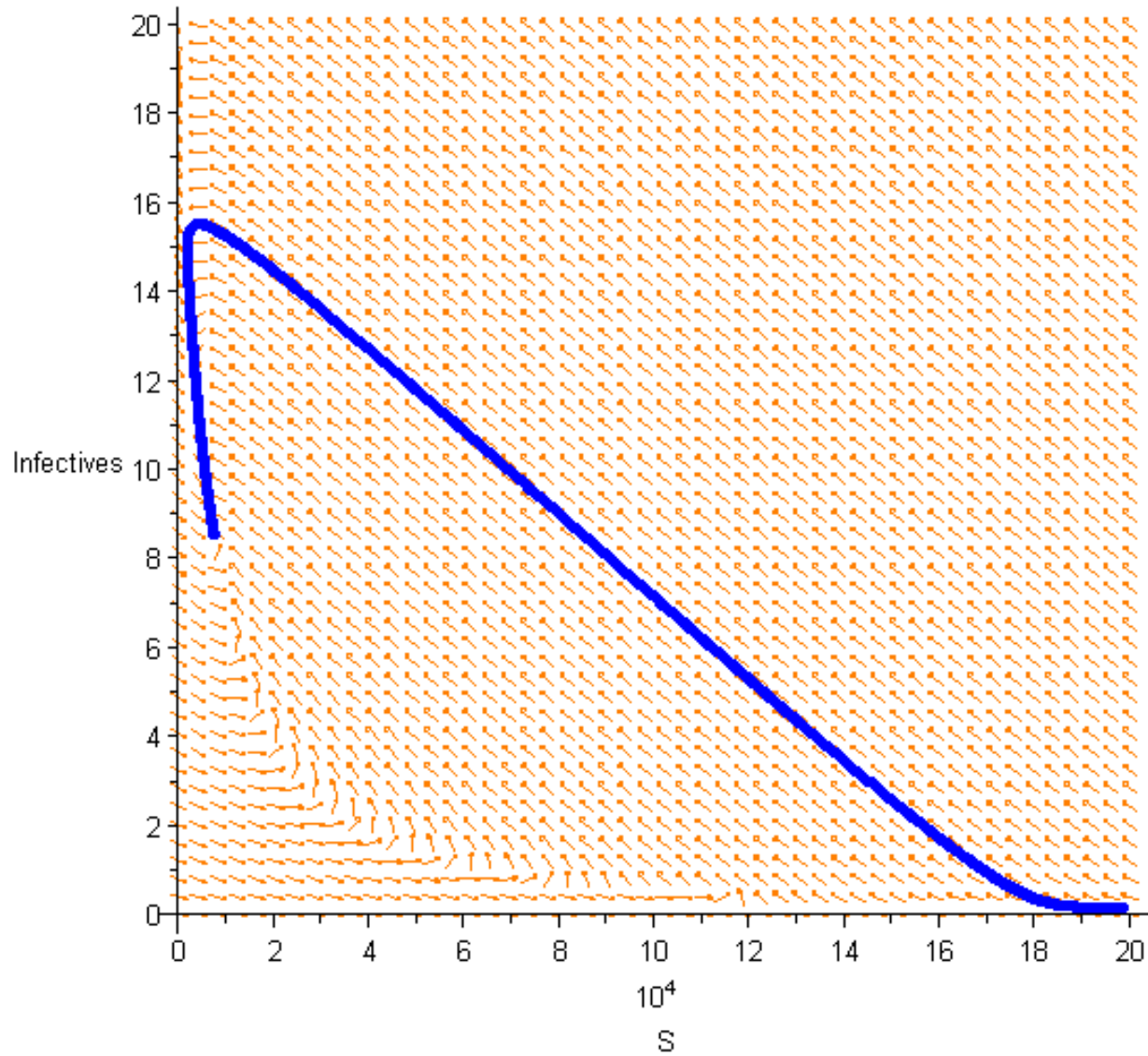
# TB In SK



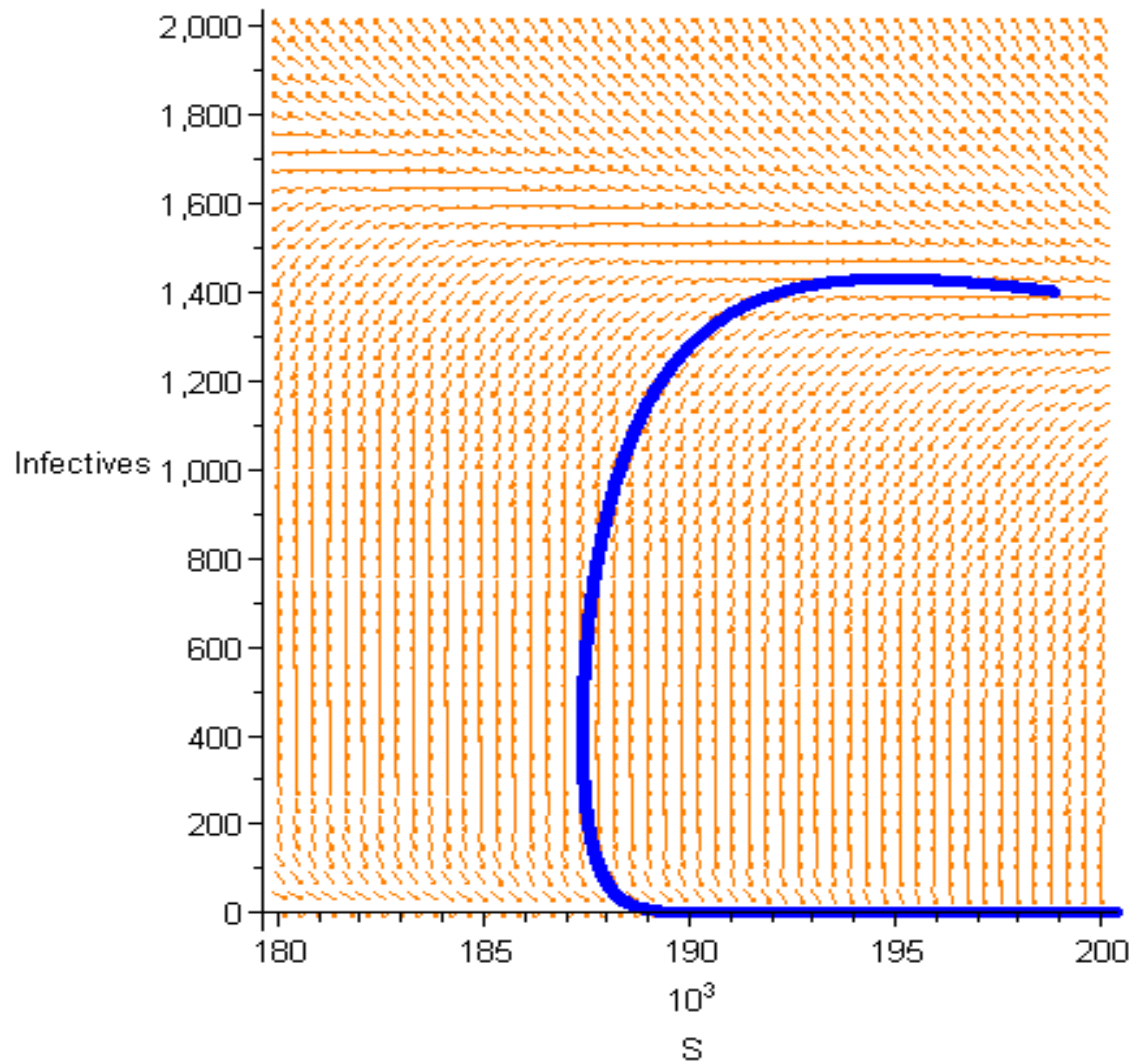
# Example: STIs



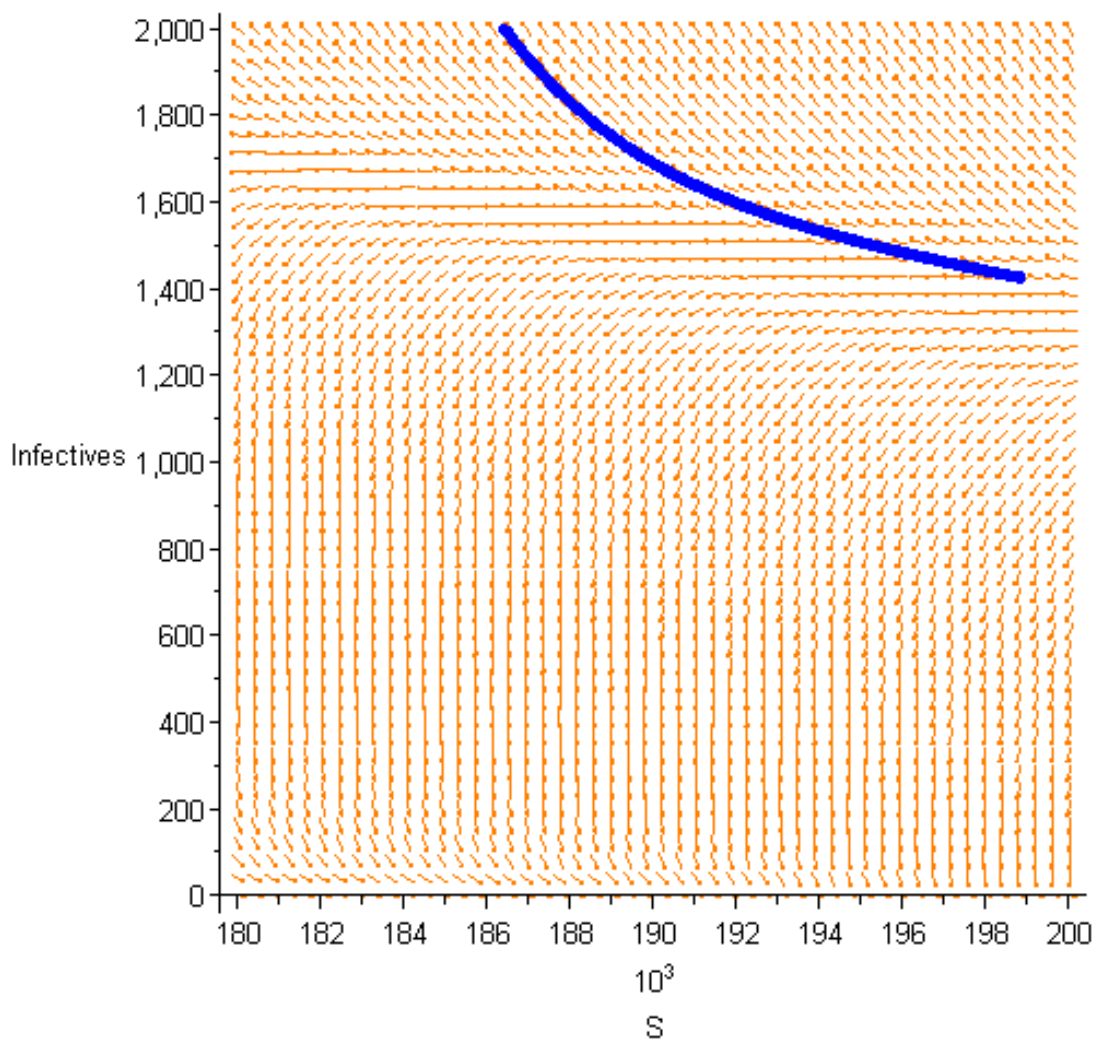
$R_0 < 1$  : 200 HC Workers,  $I_0 = 1425$



$R_0 < 1$  : 200 HC Workers,  $I_0 = 1400$

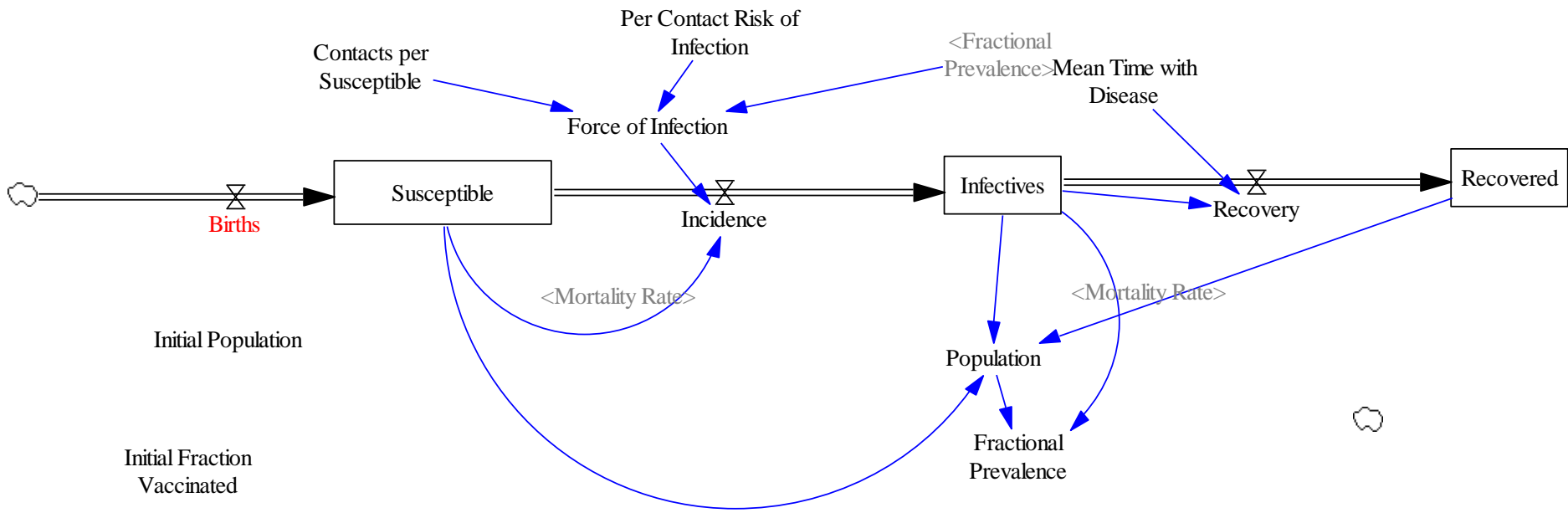


$R_0 < 1$  : 200 HC Workers,  $I_0 = 1425$



# Kendrick-McKermack Model

- Partitioning the population into 3 broad categories:
  - Susceptible (S)
  - Infectious (I)
  - Removed (R)



# Shorthand for Key Quantities for Infectious Disease Models: Stocks

- $I$  (or  $Y$ ): Total number of infectives in population
  - This could be just one stock, or the sum of many stocks in the model (e.g. the sum of separate stocks for asymptomatic infectives and symptomatic infectives)
- $N$ : Total size of population
  - This will typically be the sum of all the stocks of people
- $S$  (or  $X$ ): Number of susceptible individuals



# Key Quantities for Infectious Disease Models: Parameters

- Contacts per susceptible per unit time:  $c$ 
  - e.g. 20 contacts per month
  - This is the number of contacts a given susceptible will have with *anyone*
- Per-infective-with-susceptible-contact transmission probability:  $\beta$ 
  - This is the per-contact likelihood that the pathogen will be transmitted from an infective to a susceptible with whom they come into a single contact.

# Intuition Behind Common Terms

- $I/N$ : The Fraction of population members (or, by assumption, contacts!) that are infective
  - Important: Simplest models assume that this is also the fraction of a given susceptible's contacts that are infective! Many sophisticated models relax this assumption
- $c(I/N)$ : Number of *infectives* that come into contact with a susceptible in a given unit time
- $c(I/N)\beta$ : “Force of infection”: *Likelihood a given susceptible will be infected per unit time*
  - The idea is that if a given susceptible comes into contact with  $c(I/N)$  infectives per unit time, and if each such contact gives  $\beta$  likelihood of transmission of infection, then that susceptible has roughly a total likelihood of  $c(I/N)\beta$  of getting infected per unit time (e.g. month)

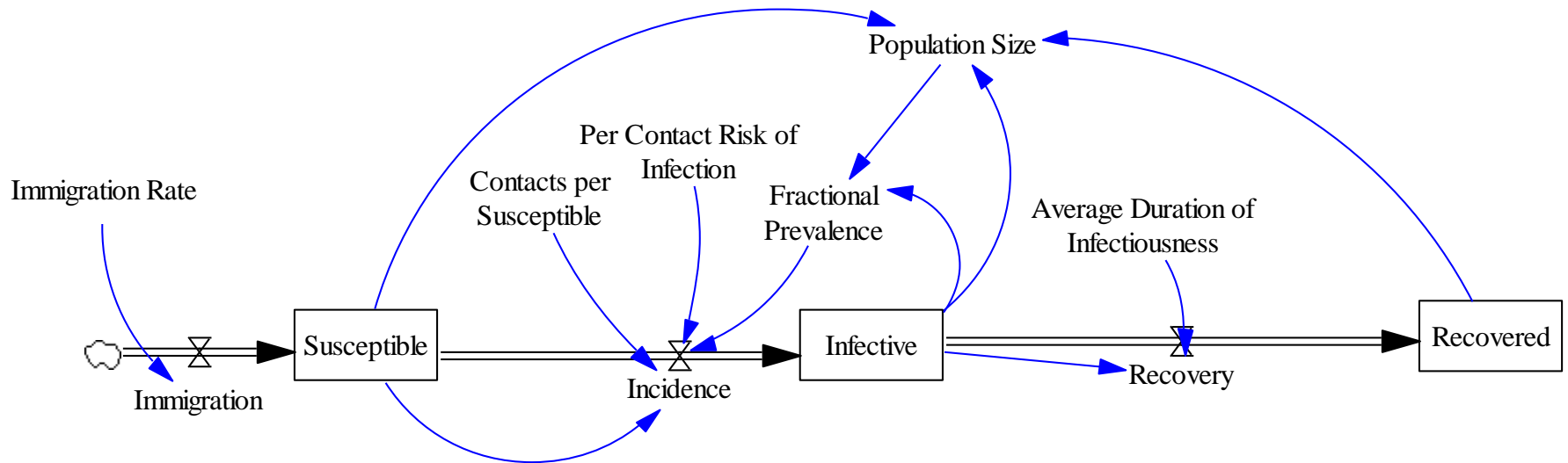
# Key Term: Flow Rate of New Infections

- This is the key form of the equation in many infectious disease models
- Total # of susceptiblesinfected per unit time  
# of Susceptibles \* “Likelihood” a given susceptible will be infected per unit time =  $S * (\textit{Force of Infection})$   
 $= S(c(I/N)\beta)$ 
  - Note that this is a term that multiplies both S and I !
    - This is much different than the purely linear terms on which we have previously focused
  - “Likelihood” is actually a likelihood density (e.g. can be  $>1$  – indicating that mean time to infection is  $<1$ )

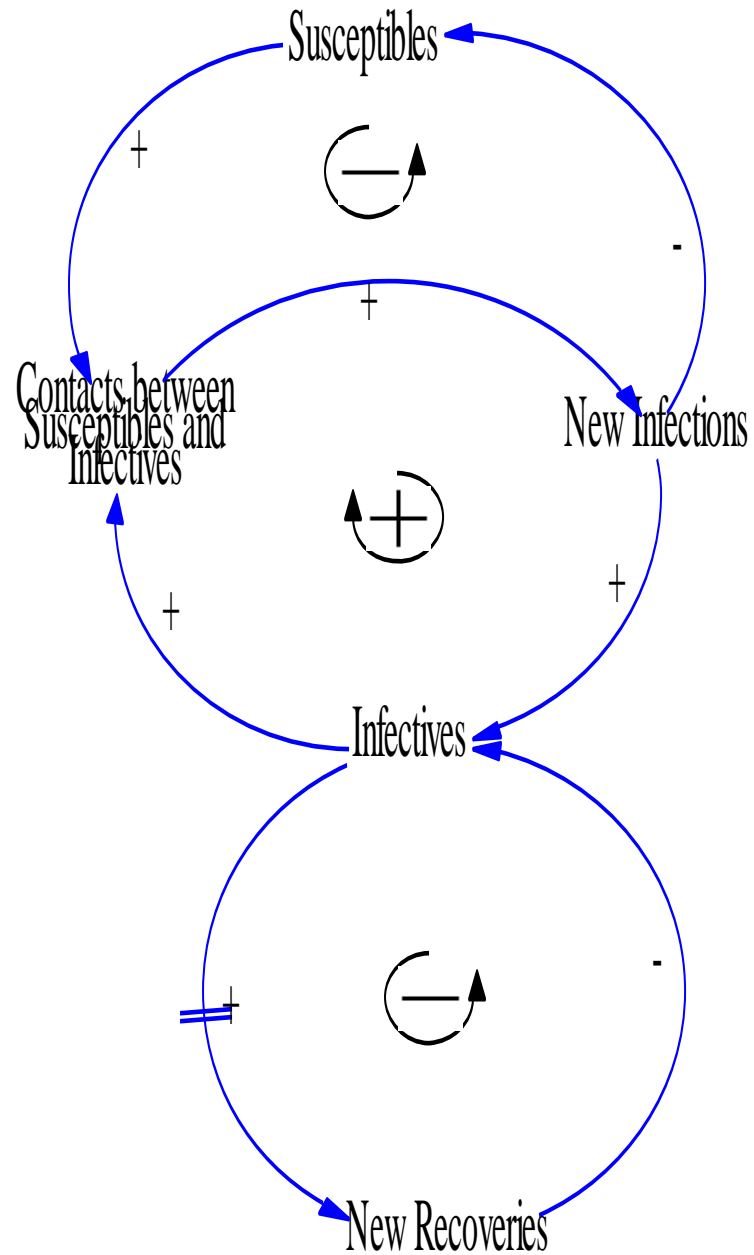
# Another Useful View of this Flow

- Recall: Total # of susceptibles infected per unit time = # of Susceptibles \* “Likelihood” a given susceptible will be infected per unit time =  $S * (\text{“Force of Infection”}) = S(c(I/N)\beta)$
- The above can also be phrased as the following:  $S(c(I/N)\beta) = I(c(S/N)\beta) = \# \text{ of Infectives} * \text{Average \# susceptibles infected per unit time by each infective}$
- This implies that as # of susceptibles falls  $\Rightarrow$  # of susceptibles surrounding each infective falls  $\Rightarrow$  the rate of new infections falls (“Less fuel for the fire” leads to a smaller burning rate)

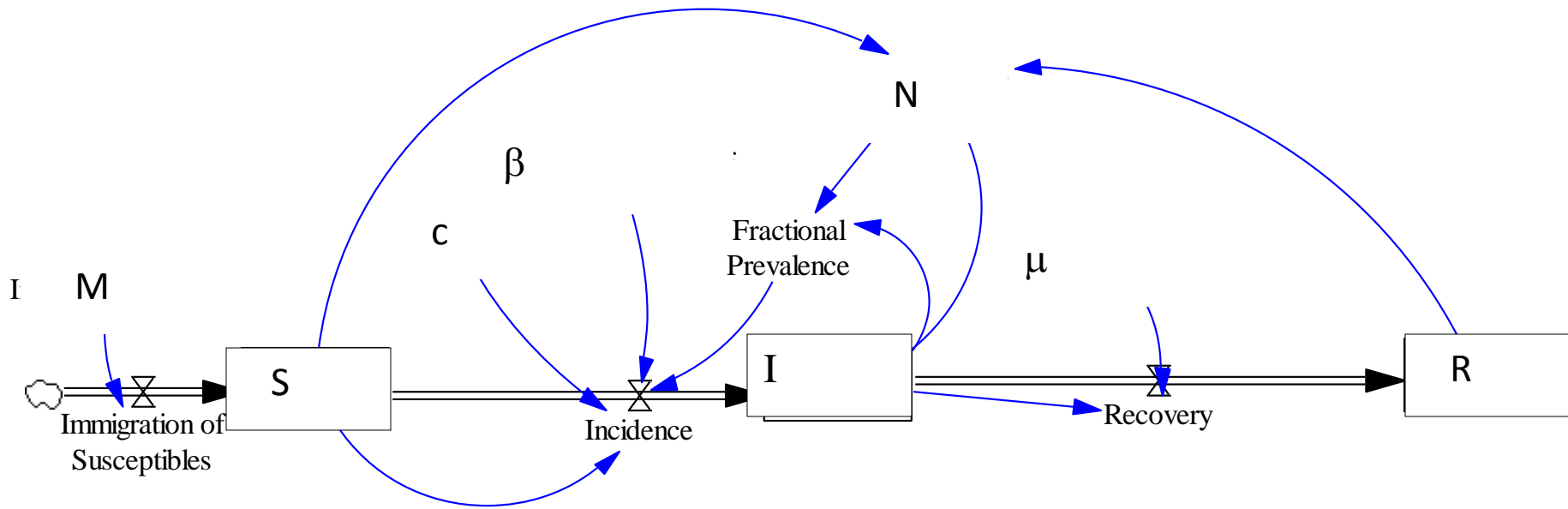
# Basic Model Structure



# Associated Feedbacks



# Mathematical Notation



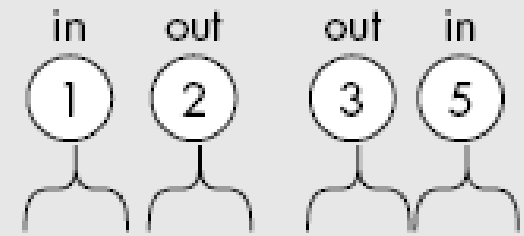
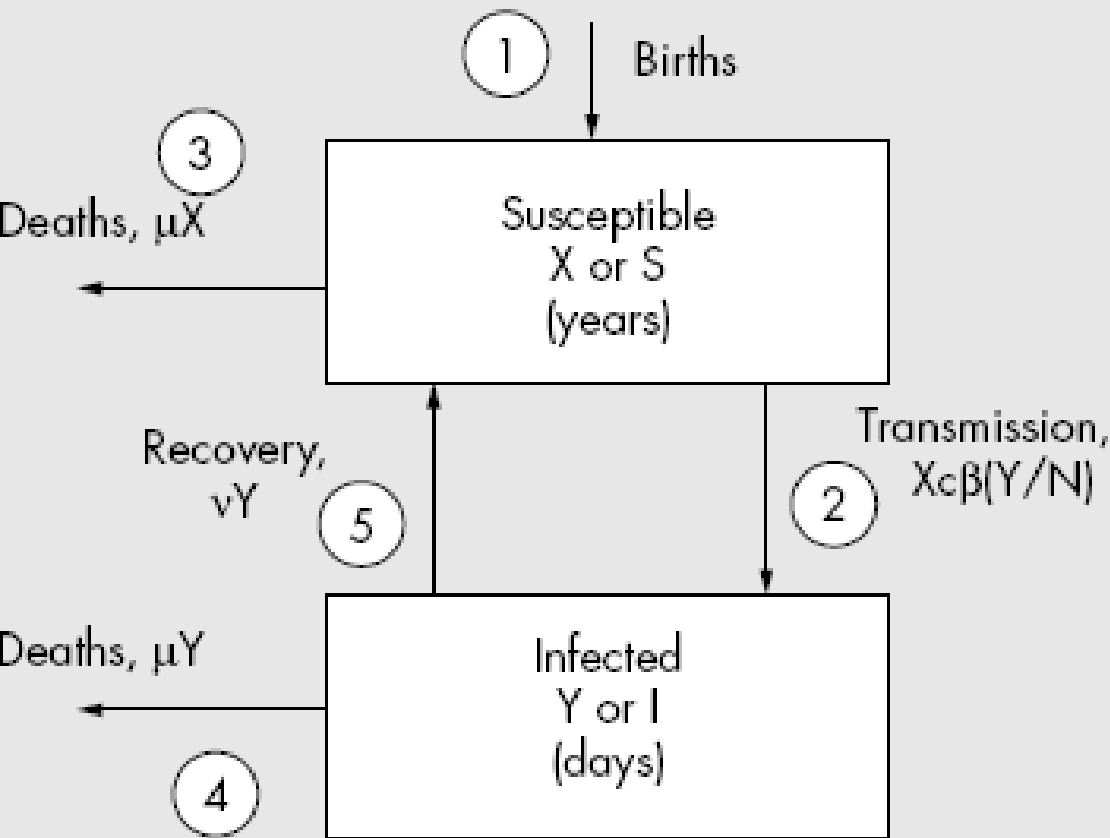
# Underlying Equations

$$\dot{S} = M - c \left( \frac{I}{N} \right) \beta S$$

$$\dot{I} = c \left( \frac{I}{N} \right) \beta S - \frac{I}{\mu}$$

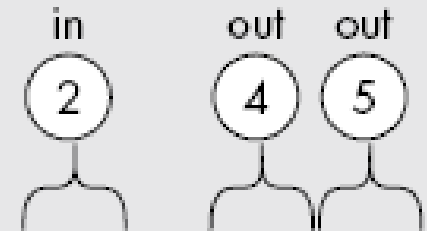
$$\dot{R} = \frac{I}{\mu}$$





$$\frac{dX}{dt} = +B - Xc\beta \frac{Y}{N} - \mu X - vY$$

+



$$\frac{dY}{dt} = +Xc\beta \frac{Y}{N} - \mu Y - vY$$

Total population  $N = X + Y$

$$\frac{dN}{dt} = +B - \mu X - \mu Y$$

Figure from Garnett, G. P. (2002). An introduction to mathematical models in sexually transmitted disease epidemiology. *Sexually Transmitted Infections*, 78(1), 7-12.